```
RESULT 1
MMII251645
                                          mRNA
                                  990 bp
                                                 linear ROD 24-NOV-2000
LOCUS
DEFINITION Mus musculus mRNA for farnesylated-proteins converting enzyme-2
           (Face-2 gene).
           AJ251645
ACCESSION
VERSION
           AJ251645.1 GI:11342585
KEYWORDS
           Face-2 gene; farnesylated-proteins converting enzyme.
SOURCE
           Mus musculus (house mouse)
 ORGANISM Mus musculus
           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
           Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
           Sciurognathi; Muroidea; Muridae; Murinae; Mus.
REFERENCE
 AUTHORS
           Cadinanos, J. and Freije, J.M.P.
 TITLE
           Characterization and expression analysis of the gene encoding the
           murine caax protease Face-2
 JOURNAL
           Unpublished
REFERENCE
           2 (bases 1 to 990)
 AUTHORS
           Freije, J.M.P.
 TITLE
           Direct Submission
           Submitted (23-NOV-1999) Freije J.M.P., Bioquimica y Biologia
 JOURNAL
           Molecular, Universidad de Oviedo, C/Julian Claveria s/n, Oviedo,
           33006, SPAIN
FEATURES
                   Location/Qualifiers
                   1. .990
    source
                   /organism="Mus musculus"
                   /mol_type="mRNA"
                   /db_xref="taxon:10090"
                   1. .990
    gene
                   /gene="Face-2"
    CDS
                   1. .990
                   /gene="Face-2"
                   /function="proteolytic processing of prenylated proteins"
                   /codon start=1
                   /product="farnesylated-proteins converting enzyme-2"
                   /protein_id="CAC17014.1"
                   /db_xref="GI:11342586"
                   /db xref="GOA:P57791"
                   /db_xref="UniProtKB/Swiss-Prot:P57791"
                   /translation="MAALGGDGLRLLSVSRPERQPESAALSGPGSGLCCWVSVFSCFS
                   LACSYVGSLYVWKSELPRDHPAVIKRRSTSVLVVSSLSPLCVLLWRELTGIQPGTSLL
                   \verb|TLMGFRLEGIFPAALLPLLLTMILFLGPLMQLSMDCPCDLTDGLKVVLAPRSWARCLT|\\
                   DMRWLRNOVIAPLTEELVFRACMLPMLAPCTGLGPAVFTCPLFFGVAHFHHIIEOLRF
                   RQSSVGSIFVSAAFQFSYTAVFGAYTAFLFIRTGHLIGPVLCHSFCNYMGFPAVCAAL
                   EHPQKWPLLAGYALGVGLFLLLLQPLTDPKLYGSLPLCMLLERTGASETLLCS"
ORIGIN
 Query Match 97.1%; Score 362; DB 6; Length 990; Best Local Similarity 99.7%; Pred. No. 4.1e-93; Matches 373; Conservative 0; Mismatches 0; Indels
                                                           1; Gaps
0v
           1 TTTGGAGTCGCCCATTTTCACCACATTATTGAGCAGCTGCGCTTCCGCCAGAGCAGTGTG 60
             610 TTTGGAGTCGCCCATTTTCACCACATTATTGAGCAGCTGCGCTTCCGCCAGAGCAGTGTG 669
          61 \ \mathsf{GGAAGTATCTTCGTGTCTGCAGCGTTCCAGTTCTCCTACACCGCTGTCTTCGGTGCTTAT \ 120
Qу
             670 GGAAGTATCTTCGTGTCTGCAGCGTTCCAGTTCTCCTACACCGCTGTCTTCGGTGCTTAT 729
         121 ACAGCTTTCCTCTTCATCCGCACAGGACACCTGATAGGGCCGGTTCTCTGCCACTCTTTC 180
Qv
             730 ACAGCTTTCCTCTTCATCCGCACAGGACACCTGATAGGGCCGGTTCTCTGCCACTCTTTC 789
         181 TGCAACTACATGGGCTTCCCTGCAGTGTGTGCAGCCCTGGAGCATCCACAGAAGTGGCCA 240
Qу
             790 TGCAACTACATGGGCTTCCCTGCAGTGTGTGCAGCCCTGGAGCATCCACAGAAGTGGCCA 849
         241 CTGCTGGCAGGCTATG-CCTCGGTGTGGGACTTTTCCTGCTTCTGCTTCAACCCCTGACA 299
Qу
             850 CTGCTGGCAGGCTATGCCCTCGGTGTGGGACTTTTCCTGCTTCTGCTTCAACCCCTGACA 909
Db
```

QУ	300	GACCCCAAGCTCTATGGCAGCCTTCCTCTTTGTATGCTTTTGGAAAGAACAGGGGCCCTCA 359
Db	910	GACCCCAAGCTCTATGGCAGCCTTCCTCTTTGTATGCTTTTGGAAAGAACAGGGGCCTCA 969
Qy	360	GAGACCCTACTGTG 373
Db	970	GAGACCCTACTGTG 983